# SOC-GA 2332 Intro to Stats Lab 8

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## 10/25/2024

# Part 1: Multivariate Regression & Interaction with One Dummy

### **Dummies**

- For categorical variables, we create dummies or convert them to 0 or 1 dummies when we want to include them in a regression model
- Note that for a categorical variable that have n categories, the regression model will only have n-1 dummies or categorical variable predictors, because the  $n^{th}$  dummy is redundant given that if an observation does not belong to any of the n-1 category, then it must be in the  $n^{th}$  category
- We call the left-out category the reference category
- Question: what if we include all n categories?
- You should always interpret your model coefficients with the reference category in mind. This could
  get complicated when you have multiple dummy variables, especially when they are interacted in your
  model

In the case of the dummies representing "race" in the earnings\_df that we will be using today, we have:

Category	$Dummy_1(black)$	$\overline{Dummy_2(other)}$
White	0	0
Black	1	0
Other	0	1

#### Exercise

- 1. Import earnings\_df.csv to your environment. Perform the following data cleaning steps: (1) If age takes the value 9999, recode it as NA; (2) Create a new variable female that equals 1 when sex takes the value female, and equals to 0 otherwise; (3) Create a new variable black that equals 1 when race is black and equals to 0 otherwise; (4) Create a new variable other that equals to 1 when race is 'other' and 0 otherwise.
- 2. Use the describe() function from the psych package to generate a quick descriptive statistics of your data.
- 3. Now, estimate the following models and display your model results in a single table using stargazer(m\_1, m\_2, ..., m\_n, type="text").
- (1) Model 1: earn  $\sim$  age (baseline)
- (2) Model 2:  $earn \sim age + edu$
- (3) Model 3:  $earn \sim age + edu + female$
- (4) Model 4:  $earn \sim age + edu + female + race$
- (5) Model 5:  $earn \sim age + edu + female + race + edu*female$

4. Write down your prediction equation for Model 5

$$I_i = \beta_0 A_i + \beta_1 E_i + \beta_2 F_i + \beta_3 B_i + \beta_4 O_i + \beta_5 E_i \times F_i$$

- 5. In Model 5, holding other variables constant, what will be the predicted difference in estimated mean earnings for a white man and a white women?
- 6. Holding other variables constant, what will be the predicted difference in estimated mean earnings for a white women and a black women?

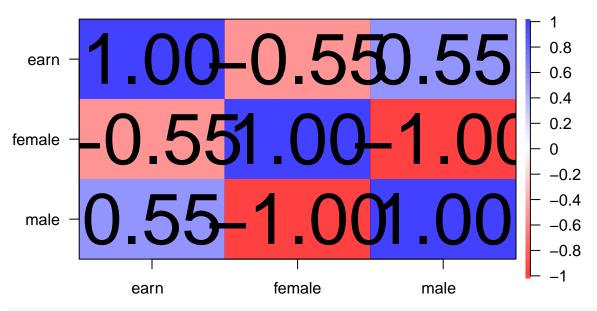
White woman = 
$$\beta_0 \bar{A}_i + \beta_1 \bar{E}_i$$
White woman =  $\beta_0 \bar{A}_i + \beta_1 \bar{E}_i + \beta_2 + \beta_5 \bar{E}_i$ 

7. Holding other variables constant, what will be the predicted difference in estimated mean earnings for a white man and a black woman?

White man =  $\beta_0 \bar{A}_i + \beta_1 \bar{E}_i$ Black woman =  $\beta_0 \bar{A}_i + \beta_1 \bar{E}_i + \beta_2 + \beta_3 + \beta_5 \bar{E}_i$ 

```
earnings_df <- read.csv("data/earnings_df.csv", stringsAsFactors = F)</pre>
## recode age
earnings_df <- earnings_df %>%
  mutate(age = case_when(
    age==9999 ~ NA,
    .default = age
  ))
## recode female
earnings_df <- earnings_df %>%
  mutate(gender = case_when(
    sex == "female" ~ 1,
    .default = 0))
## base R way of doing it
earnings_df$female <- 0</pre>
earnings_df[earnings_df$sex=="female", "female"] <- 1</pre>
## create black and other
earnings_df <-
  earnings_df %>%
  mutate(black = case_when(
    race == "black" ~ 1,
    .default = 0
  )) %>%
  mutate(other = case when(
    race == "other" ~ 1,
    .default = 0
  ))
earnings df <- earnings df %>% mutate(male = 1-female)
corPlot(earnings_df[,c("earn","female","male")])
```

# Correlation plot from data



```
##
##
                                Dependent variable:
##
##
                                        earn
##
                   (1)
                              (2)
                                        (3)
                                                   (4)
                                                               (5)
                0.134*** 0.132***
                                      0.160***
                                                 0.158*** 0.156***
##
##
                 (0.042)
                          (0.033)
                                      (0.022)
                                                 (0.022)
                                                            (0.019)
##
                                                 4.477*** 6.083***
## edu
                          4.314***
                                      4.500***
##
                           (0.171)
                                     (0.112)
                                                 (0.112)
                                                            (0.143)
##
                                     -20.528*** -20.572*** -1.571
## female
```

## ##				(0.568)	(0.565)	(1.311)
	black				-2.307*** (0.623)	
	other				-0.767 (1.137)	
## ##	edu:female					-3.128*** (0.199)
## ##	Constant				26.429*** (1.230)	
	Observations Adjusted R2		980 0.399	980 0.743	980 0.746	980 0.797
## ##	Note: *p<0.1; **p<0.05; ***p<0.07					

# Part 2: Interaction with Two Dummy Variables

Given the following modeling result, please answer the questions.

Table 1:

	Dependent variable:	
	earn	
college	6.129***	
	(0.187)	
black	-2.773***	
	(0.183)	
college:black	1.496***	
	(0.340)	
Constant	15.077***	
	(0.102)	
Observations	5,000	
$\mathbb{R}^2$	0.290	
Adjusted R <sup>2</sup>	0.289	
Residual Std. Error	5.026 (df = 4996)	
F Statistic	$679.910^{***} (df = 3; 4996)$	
Note:	*p<0.1; **p<0.05; ***p<0.0	

- 1. What will be the predicted difference in estimated mean earnings for a white person with a college degree and a black person with a college degree? Whose earnings will be higher?
- 2. What will be the predicted difference in estimated mean earnings for a white person with a college

degree and a black person without a college degree? Whose earnings will be higher?

- 3. How to interpret the interaction coefficient?
- 4. How to interpret the intercept?

#### Plot Predicted Effects

## 5

- We can visualize the predicted effects of key predictors using the predict() function in base R.
- The idea behind this task is to first create a dataframe with values of all the predictors included in the model, with only the value of your predictor(s) of interest vary within the possible range, whereas other predictors held at their mean.
- For example, if we want to examine the effect of **education and gender** on earnings, we create a dataframe with a variable **edu** that varies from 0 to 15 with an interval of 1 (so **edu** = 0, 1, 2, ..., 14, 15), because the possible value of **edu** in our data is integers from 0 to 15 (you can use **summary(your\_df)** to check value ranges).
- We repeat this number sequence for another time so that we have **each level of education for both** male and female. So we need to generate edu = 0, 1, 2, ..., 14, 15, 0, 1, 2, ..., 14, 15. We use rep(0:15, 2) to generate this number sequence.
- rep(x, times) replicate x (a vector or list) for user-defined times (in our case, times = 2). You can run this in your R console to see what number sequence is returned.
- Then, we generate a dummy variable female that equals to 0 for male and 1 for female.
- To create a dataframe that have the combination of each level of edu and each gender category, we let female = 0 for 16 times, and female = 1 for 16 times, using c(rep(0, 16), rep(1, 16)). You can run this in your R console to see what number sequence is returned.
- For the rest of the predictors, we fix them at their mean. We add na.rm = T in the mean() function to specify how we want to deal with NA values. If you don't include na.rm = T, mean() will return NA if your variable contains NAs.

```
## first, we create a dataframe with all predictor variables
## only the key predictor varies, while the others remain at the mean
pred_IV <- data.frame(edu = rep(0:15, 2)) %>%
                                                     ## first, create a df with values of your key pre
  mutate(female = c(rep(0, 16), rep(1, 16)),
                                                     ## b/c we are looking at interaction effects,
                                                     ## give gender two values, otherwise fix it at me
         age = mean(earnings_df$age, na.rm = T),
                                                  ## fix other variables at mean
         black = mean(earnings_df$black),
         other = mean(earnings_df$other))
rep(0:15,2)
## [1]
        0 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 0 1 2 3 4 5
## [26]
        9 10 11 12 13 14 15
## examine the df
head(pred_IV, 5)
     edu female
                     age black other
## 1
      0
             0 43.26429 0.306 0.068
## 2
             0 43.26429 0.306 0.068
      1
             0 43.26429 0.306 0.068
## 3
      2
## 4
       3
              0 43.26429 0.306 0.068
```

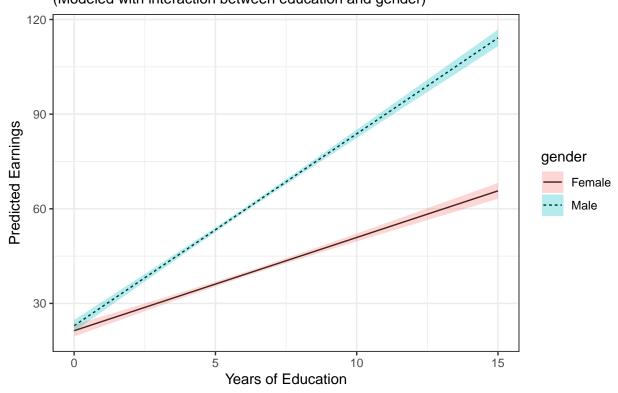
• Now that we have the dataframe pred\_IV ready for predicting the dependent variable (earning), we can use the R function predict() to calculate fitted earning using the regression model and the values

0 43.26429 0.306 0.068

specified in each row in pred\_IV. Then, use cbind() to combine this fitted Y value vector with your pred\_IV for plotting.

```
## use `predict` to predict the Y
predicted_earning <- predict(m5,</pre>
                                                       ## the model you are using
                             pred_IV,
                                                      ## the df you use for predicting
                             interval = "confidence", ## set CI
                             level = 0.95)
## bind the columns
pred_result <- cbind(pred_IV, predicted_earning)</pre>
## check df
head(pred_result, 5)
     edu female
                     age black other
                                          fit
                                                    lwr
## 1
              0 43.26429 0.306 0.068 22.93127 21.12317 24.73936
              0 43.26429 0.306 0.068 29.01392 27.46140 30.56644
## 3
              0 43.26429 0.306 0.068 35.09657 33.78940 36.40374
## 4
       3
              0 43.26429 0.306 0.068 41.17922 40.10017 42.25827
## 5
              0 43.26429 0.306 0.068 47.26188 46.38025 48.14350
## plot
pred_result %>%
  mutate(gender = ifelse(female == 0, "Male", "Female")) %>%
                                                                    ## convert dummy to character variab
  ggplot(aes(x = edu, y = fit, group = gender)) +
  geom_line(aes(linetype = gender)) +
                                                                    ## group linetype by gender
  geom_ribbon(aes(ymin = lwr, ymax = upr, fill = gender), alpha = 0.3) + # add 95% CI
  theme bw() +
  labs(x = "Years of Education",
       y = "Predicted Earnings") +
  ggtitle("Predicted Earnings by Education and Gender",
          subtitle = "(Modeled with interaction between education and gender)")
```

# Predicted Earnings by Education and Gender (Modeled with interaction between education and gender)



## Part 3 F-test for Nested Models

- We can use F-test to compare two regression models. The idea behind the F-test for nested models is to check **how much errors are reduced after adding additional predictors**. A relatively large reduction in error yields a large F-test statistic and a small P-value. The P-value for F statistics is the right-tail probability.
- If the F's p-value is significant (smaller than 0.05 for most social science studies), it means that at least one of the additional  $\beta_i$  in the full model is not equal to zero.
- The F test statistic for nested regression models is calculated by:

$$F = \frac{(SSE_{\text{restricted}} - SSE_{\text{full}})/df_1}{SSE_{\text{full}}/df_2}$$

where  $df_1$  is the number of **additional** predictors added in the full model and  $df_2$  is the **residual df for the** full **model**, which equals (n-1) number of IVs in the complete model). The df of the F test statistic is  $(df_1, df_2)$ .

For example, according to the equation, we can hand-calculate the F value for m3 vs m4:

```
# SSE_restricted:
sse_m3 <- sum(m3$residuals^2)

# SSE_full:
sse_m4 <- sum(m4$residuals^2)

# We add one additional IV, so:</pre>
```

```
# Residual df for the full model (m5):
df2 <- m4$df.residual

# Calculate F:
F_stats <- ((sse_m3 - sse_m4)/df1)/(sse_m4/df2)
F_stats

## [1] 6.855912

# Check tail probability using `1 - pf()`
1 - pf(F_stats, df1, df2)</pre>
```

## [1] 0.001104788

• You can also use anova() to perform a F-test in R.

```
anova(m3, m4)
```

```
## Analysis of Variance Table
##
## Model 1: earn ~ age + edu + female
## Model 2: earn ~ age + edu + female + black + other
              RSS Df Sum of Sq
     Res.Df
                                         Pr(>F)
##
                                     F
## 1
        976 76776
## 2
        974 75711
                        1065.8 6.8559 0.001105 **
## ---
                  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
```

• What is your null and alternative hypotheses? What's your decision given the F-test result?

# Part 4: Standardized Regression Coefficients

- Why sometimes people report standardized regression coefficients? As we covered in the lecture, the size of a regression coefficient depends on the scale at which the independent and dependent variables are measured.
- For example, assume that in a regression model the coefficient of population on the national GDP is 0.0001. This means that 1 additional person will lead to 0.0001 increase in the GDP. However, this value does not necessarily imply that the effect of population is less pronounced than other predictors whose coefficients have a larger value. Because the value of the coefficient depends on the measurement unit of the IV. If we now change population to **population in million**, the new coefficient of population will become  $0.0001 \cdot 10^6 = 100$ . Although the value of the coefficient gets much larger, this increase is caused by a change in the measurement unit, not the actual effect of population.
- Therefore, it is problematic to use the raw value of the regression coefficient as indicators of relative effect size if the predictors in the model have different measurement units. In such scenarios, standardized regression coefficients can help compare the relative effect size of the predictors even if they are measured in different units.
- Standardized coefficients convert both your dependent variable and independent variables to **z-scores**. That is, each of your original (numeric) variables are converted to have a mean of 0 and a standard deviation of 1. Thus, standardized coefficients tell us the change in Y, in Y's standard deviation units, for a one-standard-deviation increase in  $X_i$ , while holding other Xs constant.
- There are two methods of getting standardized regression coefficients in R.

## Method 1: Use lm.beta() from the QuantPsyc package

You can get standardized regression coefficients by using the lm.beta() function in the QuantPsyc package. For example, if we want to get the standardized coefficients for Model 2 (earn ~ age\_recode + edu):

```
original model
m2
##
## lm(formula = earn ~ age + edu, data = earnings_df)
## Coefficients:
## (Intercept)
                                      edu
                         age
       17.7857
##
                      0.1321
                                   4.3140
## standardized coefficients
std m2 <- lm.beta(m2)
std_m2
##
                      edu
          age
## 0.09914669 0.62457139
```

• But this method will only report the point estimates instead of a comprehensive modeling result. To obtain that, we need to convert all numeric variables to z-scores and estimate regression models based on the transformed data.

### Method 2: Create Z-scores for All Numeric Variables

##

• For each numeric variables, we create the "standardized variables" by calculating their z-scores:

$$z = \frac{x - \bar{x}}{s_x}$$

• For example, we can use mutate\_at() to covert numeric variables to z-scores in earnings\_df using the above formula:

```
## a function that convert a numeric vector to a z-score vector
get_zscore <- function(x){</pre>
  (x - mean(x, na.rm = T))/sd(x, na.rm = T)
  }
## create a df with numeric variables converted to z-score
earnings_df_std <- earnings_df %>%
  mutate_at(c("edu", "age", "earn"), get_zscore)
## estimate model
m2 std zscore <- lm(earn ~ age + edu, data = earnings df std)
## compare results
stargazer(m2, m2_std_zscore, type = "text")
##
##
                                       Dependent variable:
##
##
                                               earn
```

(1)

(2)

##					
##	age	0.132***	0.098***		
##		(0.033)	(0.025)		
##					
##	edu	4.314***	0.622***		
##		(0.171)	(0.025)		
##					
##	Constant	17.786***	0.002		
##		(1.817)	(0.025)		
##					
##					
##	Observations	980	980		
##	R2	0.400	0.400		
##	Adjusted R2	0.399	0.399		
##	Residual Std. Error (df = 977)	13.558	0.770		
##	F Statistic (df = 2; 977)	326.017***	326.017***		
##					
##	Note:	*p<0.1; **p<0.	.05; ***p<0.01		